

# SEQUENCE LISTING

<110> Crane, Edmund H.  
Crane, Virginia C.

<120> Maize Pathogenesis-Related  
Polynucleotide and Methods of Use

<130> 35718/214291

<150> US 60/195,801

<151> 2000-04-10

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<221> CDS

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ccc atg gcc acg gcg tgc ttg ctc ctc gcc acc ctc ctc gcg ctc tgc	155
Pro Met Ala Thr Ala Cys Leu Leu Leu Ala Thr Leu Leu Ala Leu Cys	
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Ala Ala Pro Ala Pro Thr His Gly Ala Arg Val Leu Met Pro Gly Gly	
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gcg ggc gcg gtg acc aag gcg cag cag ggt ggc acc ggc agc ggc agc	251
Ala Gly Ala Val Thr Lys Ala Gln Gln Gly Gly Thr Gly Ser Gly Ser	
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aac gcg acg gcg gac gag tac ctg gcg ccg cac aac cag gcg cgc gcg	299
Asn Ala Thr Ala Asp Glu Tyr Leu Ala Pro His Asn Gln Ala Arg Ala	
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gcg gtg ggc gtg gcc ccg ctg cgg tgg aac gcg ggc ctg gct tcg gcg	347
Ala Val Gly Val Ala Pro Leu Arg Trp Asn Ala Gly Leu Ala Ser Ala	
80 85 90 95	

gcc gcg ggg acg gtg gcg cag cag ccg ccg cag ggc ggg tgc gcg ttc	395
Ala Ala Gly Thr Val Ala Gln Gln Arg Arg Gln Gly Gly Cys Ala Phe	
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Ala Asp Val Gly Ala Ser Pro Tyr Gly Ala Asn Gln Gly Trp Ala Ser  
115 120 125

tac cgc gcg cgc ccc gcc gag gtg gtg gcg ctg tgg gtg gcg gag ggg 491  
Tyr Arg Ala Arg Pro Ala Glu Val Val Ala Leu Trp Val Ala Glu Gly  
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cgg tac tac acc cac gcc aac aac acg tgc gcc gcg ggg cgg cag tgc 539  
Arg Tyr Tyr Thr His Ala Asn Asn Thr Cys Ala Ala Gly Arg Gln Cys  
145 150 155

ggc acg tac acg cag gtg gtg tgg cgc aac acc gcc gag gtc ggg tgc 587  
Gly Thr Tyr Thr Gln Val Val Trp Arg Asn Thr Ala Glu Val Gly Cys  
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Ala Gln Ala Ser Cys Ala Thr Gly Ala Thr Leu Thr Leu Cys Leu Tyr  
180 185 190

aac ccg cac ggc aac gtg cag ggc cag agc ccc tac tag ctagctgagg 684  
Asn Pro His Gly Asn Val Gln Gly Gln Ser Pro Tyr \*  
195 200

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35 40 45  
Gly Ala Val Thr Lys Ala Gln Gln Gly Gly Thr Gly Ser Gly Ser Asn  
50 55 60  
Ala Thr Ala Asp Glu Tyr Leu Ala Pro His Asn Gln Ala Arg Ala Ala  
65 70 75 80  
Val Gly Val Ala Pro Leu Arg Trp Asn Ala Gly Leu Ala Ser Ala Ala  
85 90 95  
Ala Gly Thr Val Ala Gln Gln Arg Arg Gln Gly Gly Cys Ala Phe Ala  
100 105 110  
Asp Val Gly Ala Ser Pro Tyr Gly Ala Asn Gln Gly Trp Ala Ser Tyr  
115 120 125  
Arg Ala Arg Pro Ala Glu Val Val Ala Leu Trp Val Ala Glu Gly Arg  
130 135 140  
Tyr Tyr Thr His Ala Asn Asn Thr Cys Ala Ala Gly Arg Gln Cys Gly  
145 150 155 160  
Thr Tyr Thr Gln Val Val Trp Arg Asn Thr Ala Glu Val Gly Cys Ala  
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Met Ala Thr Ala Cys Leu Leu Leu Ala Thr Leu Leu Ala Leu Cys Ala  
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Ala Pro Ala Pro Thr His Gly Ala Arg Val Leu Met Pro Gly Gly Ala  
35 40 45  
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Gly Ala Val Thr Lys Ala Gln Gln Gly Gly Thr Gly Ser Gly Ser Asn  
50 55 60  
gcg acg gcg gac gag tac ctg gcg ccg cac aac cag gcg cgc gcg gcg 240  
Ala Thr Ala Asp Glu Tyr Leu Ala Pro His Asn Gln Ala Arg Ala Ala  
65 70 75 80  
gtg ggc gtg gcc ccg ctg cgg tgg aac gcg ggc ctg gct tcg gcg gcc 288  
Val Gly Val Ala Pro Leu Arg Trp Asn Ala Gly Leu Ala Ser Ala Ala  
85 90 95  
gcg ggg acg gtg gcg cag cag cgg cgg cag ggc ggg tgc gcg ttc gcg 336  
Ala Gly Thr Val Ala Gln Gln Arg Arg Gln Gly Gly Cys Ala Phe Ala  
100 105 110  
gac gtg ggg gcc agc ccc tac ggc gcg aac cag ggg tgg gcg agc tac 384  
Asp Val Gly Ala Ser Pro Tyr Gly Ala Asn Gln Gly Trp Ala Ser Tyr  
115 120 125  
cgc gcg cgc ccc gcc gag gtg gtg gcg ctg tgg gtg gcg gag ggg cgg 432  
Arg Ala Arg Pro Ala Glu Val Val Ala Leu Trp Val Ala Glu Gly Arg  
130 135 140  
tac tac acc cac gcc aac aac acg tgc gcc gcg ggg cgg cag tgc gcc 480  
Tyr Tyr Thr His Ala Asn Asn Thr Cys Ala Ala Gly Arg Gln Cys Gly  
145 150 155 160  
acg tac acg cag gtg gtg tgg cgc aac acc gcc gag gtc ggg tgc gcg 528  
Thr Tyr Thr Gln Val Val Trp Arg Asn Thr Ala Glu Val Gly Cys Ala

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Gln Ala Ser Cys Ala Thr Gly Ala Thr Leu Thr Leu Cys Leu Tyr Asn						
	180		185		190	
ccg cac ggc aac gtg cag ggc cag agc ccc tac tag						612
Pro His Gly Asn Val Gln Gly Gln Ser Pro Tyr *						
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